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Sequence Listing was accepted.

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Reviewer: Keisha Douglas

Timestamp: [year=2008; month=10; day=27; hr=16; min=19; sec=49; ms=252;
]

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Application No: 10514626 Version No: 2.0

Input Set:**Output Set:**

Started: 2008-09-25 15:54:27.253
Finished: 2008-09-25 15:54:29.922
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 669 ms
Total Warnings: 50
Total Errors: 0
No. of SeqIDs Defined: 53
Actual SeqID Count: 53

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (21)
W 402	Undefined organism found in <213> in SEQ ID (22)

Input Set:

Output Set:

Started: 2008-09-25 15:54:27.253

Finished: 2008-09-25 15:54:29.922

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 669 ms

Total Warnings: 50

Total Errors: 0

No. of SeqIDs Defined: 53

Actual SeqID Count: 53

Error code	Error Description
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This error has occurred more than 20 times, will not be displayed	

SEQUENCE LISTING

<110> Aarhus Universitet
Pedersen, Finn Skou
Bahrami, Shervin
Duch, Mogens Ryttergaard

<120> A purified retroviral envelope polypeptide

<130> P864US00

<140> 10514626

<141> 2005-06-23

<160> 53

<170> PatentIn version 3.4

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<211> 1920

<212> DNA

<213> Murine leukemia virus

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<211> 639

<212> PRT

<213> Murine leukemia virus

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 20 25 30

Val Gln His Asp Ser Pro His Gln Val Phe Asn Val Thr Trp Arg Val
 35 40 45

Thr Asn Leu Met Thr Gly Gln Thr Ala Asn Ala Thr Ser Leu Leu Gly
 50 55 60

Thr Met Thr Asp Ala Phe Pro Lys Leu Tyr Phe Asp Leu Cys Asp Leu

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Gly Arg Lys Arg Ala Arg Ile Phe Asp Phe Tyr Val Cys Pro Gly His						
	100		105		110	
Thr Val Leu Ala Gly Cys Gly Gly Pro Arg Glu Gly Tyr Cys Gly Lys						
	115		120		125	
Trp Gly Cys Glu Thr Thr Gly Gln Ala Tyr Trp Lys Pro Ser Ser Ser						
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Trp Asp Leu Ile Ser Leu Lys Arg Gly Asn Thr Pro Lys Gly Gln Gly						
	145		150		155	160
Pro Cys Tyr Asp Ser Ser Val Val Ser Ser Ser Ala Gln Gly Ala Thr						
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Pro Gly Gly Arg Cys Asn Pro Leu Val Leu Glu Phe Thr Asp Ala Gly						
	180		185		190	
Lys Arg Ala Ser Trp Asp Ala Ser Lys Ala Trp Gly Leu Arg Leu Tyr						
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Arg Ser Thr Arg Thr Asp Pro Val Thr Arg Phe Ser Leu Thr Arg Gln						
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Val Leu Asn Ile Gly Pro Arg Val Pro Ile Gly Pro Asn Pro Val Ile						
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Ile Asp Gln Leu Pro Pro Ser Arg Pro Val Gln Ile Met Leu Pro Arg						
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Pro Pro Gln Pro Pro Pro Pro Gly Ala Ala Ser Thr Val Pro Glu Thr						
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Ala Pro Pro Ser Gln Gln Pro Gly Thr Gly Asp Arg Leu Leu Asn Leu						
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Val Asn Gly Ala Tyr Gln Ala Leu Asn Leu Thr Ser Pro Asp Lys Thr						
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Gln Glu Cys Trp Leu Cys Leu Val Ala Gly Pro Pro Tyr Tyr Glu Gly
305 310 315 320

Val Ala Val Leu Gly Thr Tyr Ser Asn His Thr Ser Ala Pro Ala Asn
325 330 335

Cys Ser Val Ala Ser Gln His Lys Leu Thr Leu Ser Glu Val Thr Gly
340 345 350

Gln Gly Leu Cys Val Gly Ala Val Pro Lys Thr His Gln Ala Leu Cys
355 360 365

Asn Thr Thr Gln Lys Thr Ser Asn Gly Ser Tyr Tyr Leu Ala Ala Pro
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Ala Gly Thr Ile Trp Ala Cys Asn Thr Gly Leu Thr Pro Cys Leu Ser
385 390 395 400

Thr Thr Val Leu Asp Leu Thr Thr Asp Tyr Cys Val Leu Val Glu Leu
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Trp Pro Lys Val Thr Tyr His Ser Pro Gly Tyr Val Tyr Gly Gln Phe
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Glu Glu Lys Thr Lys Tyr Lys Arg Glu Pro Val Ser Leu Thr Leu Ala
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Leu Leu Leu Gly Gly Leu Thr Met Gly Gly Ile Ala Ala Gly Val Gly
450 455 460

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465 470 475 480

Ala Ala Met Gln Asp Asp Leu Lys Glu Val Glu Lys Ser Ile Thr Asn
485 490 495

Leu Glu Arg Ser Leu Thr Ser Leu Ser Glu Val Val Leu Gln Asn Arg
500 505 510

Arg Gly Leu Asp Leu Leu Phe Leu Lys Glu Gly Gly Leu Cys Ala Ala
515 520 525

Leu Lys Glu Glu Cys Cys Phe Tyr Ala Asp His Thr Gly Leu Val Arg
530 535 540

Asp Ser Met Ala Lys Leu Arg Glu Arg Leu Ser Gln Arg Gln Lys Leu
545 550 555 560

Phe Glu Ser Gln Gln Gly Trp Phe Glu Gly Leu Phe Asn Lys Ser Pro
565 570 575

Trp Phe Thr Thr Leu Ile Ser Thr Ile Met Gly Pro Leu Ile Ile Leu
580 585 590

Leu Leu Ile Leu Leu Phe Gly Pro Cys Ile Leu Asn His Leu Val Gln
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Phe Ile Lys Asp Arg Val Ser Val Val Gln Ala Leu Val Leu Thr Gln
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<220>
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<223> Primer

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<211> 17

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Pro

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<212> PRT

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1 5 10 15

Pro

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